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FIG. 1A



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TNIK	779	VKPPEISRDITRPSRPSYKKAIDEDLTALAKELRIEETNRPWIKVTDYSSSESSSEEEEDGESETHDGTVAVS
NIK	716	KKPDDKKEVFR-----SLKPAGEVDTLAKELRAV--EDVRPPIKVTDLAKELRAV--
TNIK	859	WIPRLIPTGAPGSNEQYNVGMNGTHGLETSHADSFSGSISREGTLMIRETSGEKKRSGHSDSNGFAGHINLPDLVQQSSES
NIK	788	EDTRAASSSPNLSNGETESVKTWVHDDESEP--ANTIPSKKEGTLMVQTQSASS--
TNIK	939	PAGTPTEGLGRVSTHSQEMDSGTEYGMGSSTKASIFTPFIIDPRVYQTSPIDEDEEDESSAAAIFTSELLRQEQAQKLNEAR
NIK	846	-SSSSSFTPFIIDPRLQISPSSS--GTTVTSVVGESCDRGLRPEAIRQDPTR
TNIK	1019	KISVVNNVNPNTNIRPHISDTPEIRKYKKRFNSEILCAALWGVNLVGTEENGLMILDRSQGQGKVVNLINRRRFQQMDVLEGLN
NIK	892	KGSVVNNVNPNTNTRPQSDTPEIRKYKKRFNSEILCAALWGVNLVGTEESGLMILDRSQGQGKVVPLISRRRFQQMDVLEGLN
TNIK	1099	VLVTISGKKNKLRVYYLSWLRLNRILHNDPEVEKKQGWITVGDLEGCHYKVVKYERIKFLVIALKNAVEIYAWAPKPYHK
NIK	972	VLVTISGKKDKKLRLRVYYLSWLRLNRILHNDPEVEKKQGWITVGDLEGCHYKVVKYERIKFLVIALKSSVENIYAWAPKPYHK
TNIK	1179	FMAFKSFADLOHKPLLVDLTVEEGQRLKVIIFGSHITGFHVIDVDSGNSYDIYIPSHIQGNITPHAIIVLPKTDGMEMLVCY
NIK	1052	FMAFKSFGEELLHKPLLVDLTVEEGQRLKVIIFGSCAGFHAVIDVDSGSVYDIYIPSHIQCSIKPHAIIVLPNTDGMEMLVCY
TNIK	1259	EDEGVVYNTYGRITKDVVLQWGEMPTSVAIHSNQIMGWGEKAIEIERSVETGHLQDGVMFHKRAQRLKFLCERNDKVFAS
NIK	1132	EDEGVVYNTYGRITKDVVLQWGEMPTSVAIHSNQIMGWGEKAIEIERSVETGHLQDGVMFHKRAQRLKFLCGRNDKVFAS
TNIK	1339	VRSGGSSQVFFMFTLNRMNMINW
NIK	1212	VRSGGSSQVFFMFTLGRTSILSW

FIG.-1B



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FIG._2

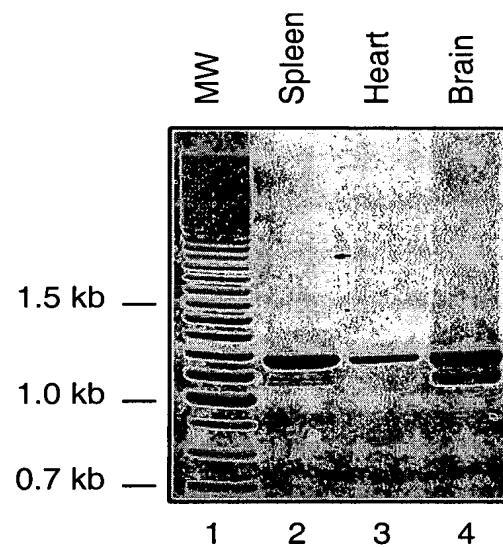


FIG._3

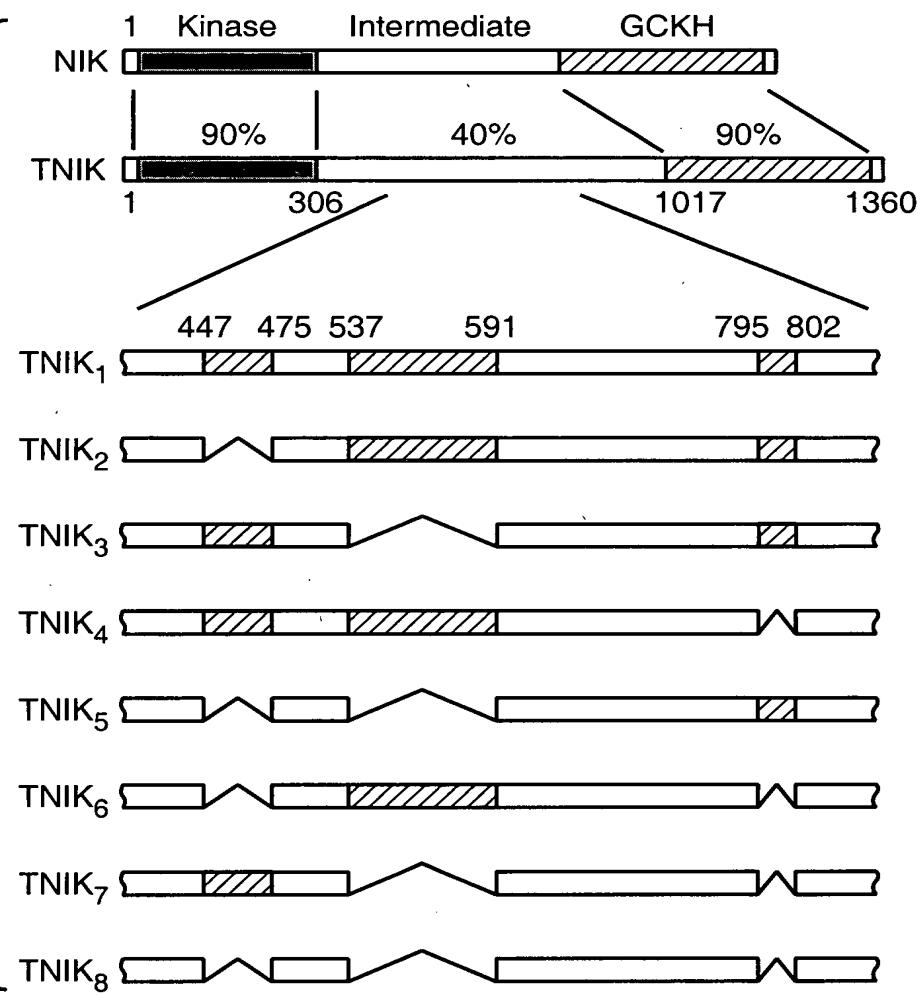




FIG.-4

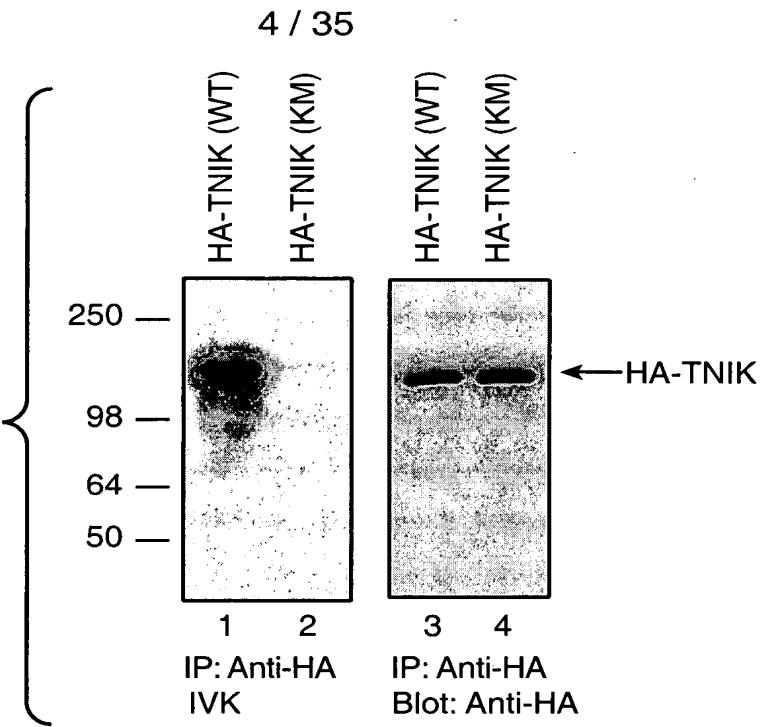


FIG.-5A

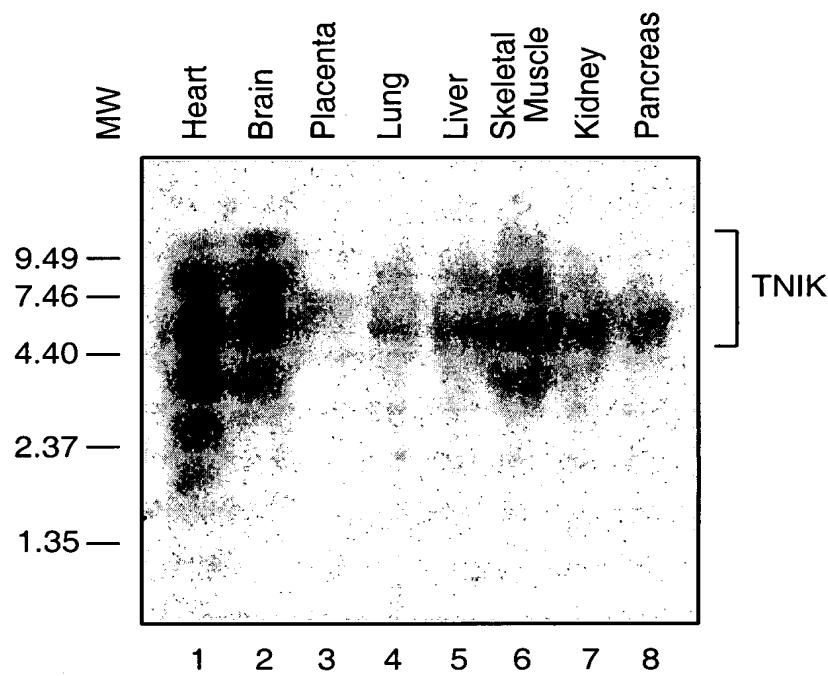
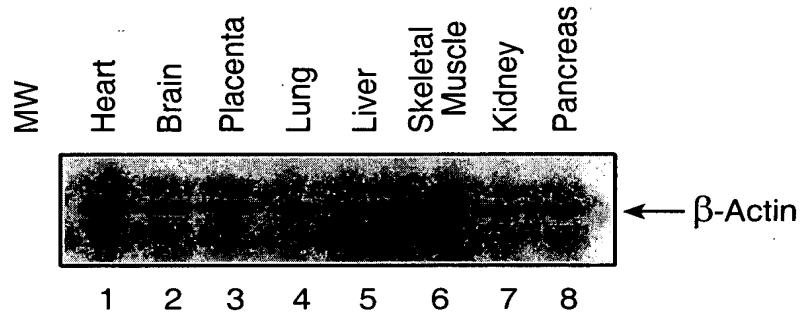
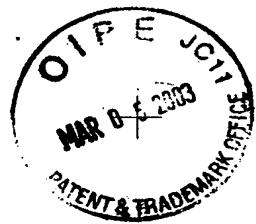


FIG.-5B





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FIG.-6

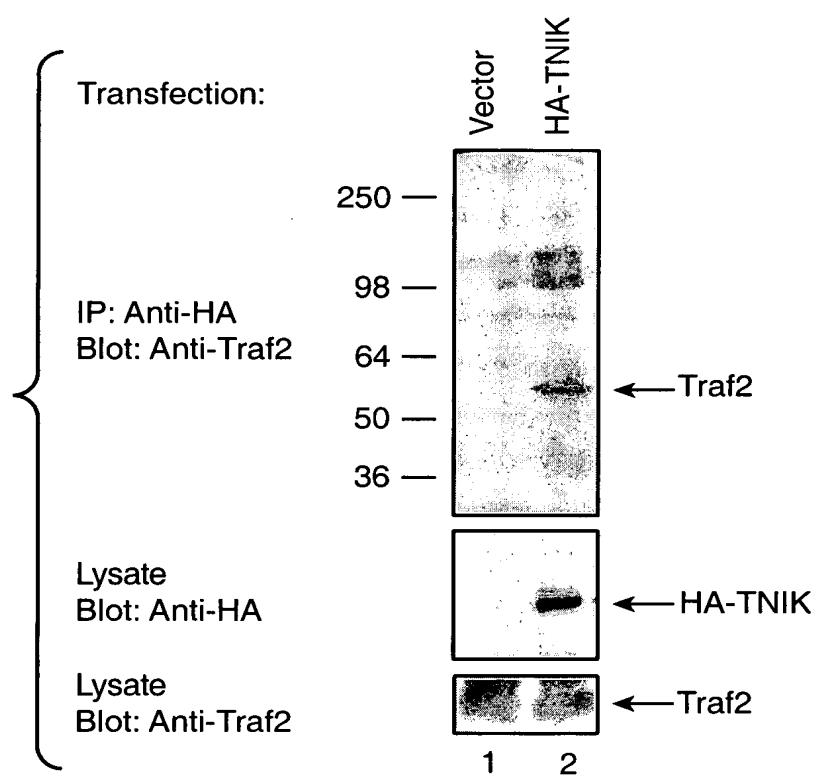
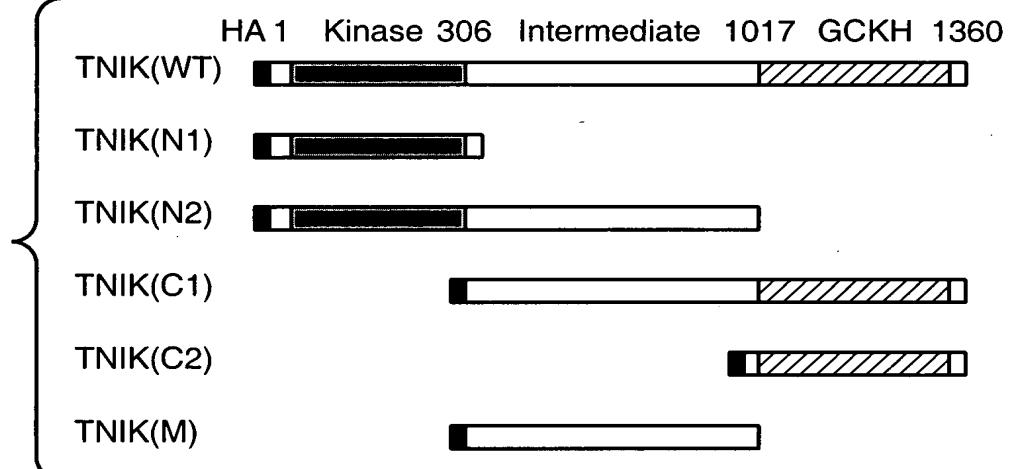


FIG.-7



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FIG._8A

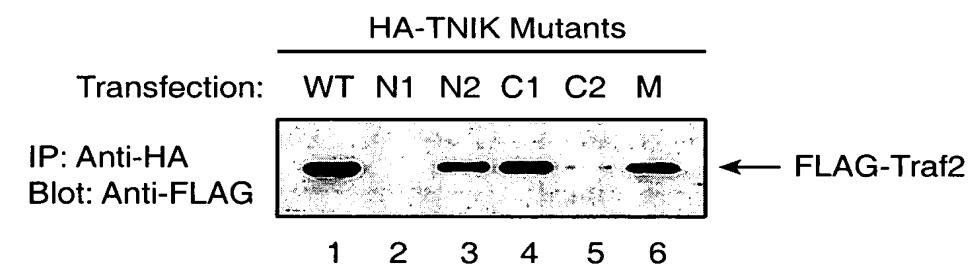


FIG._8B

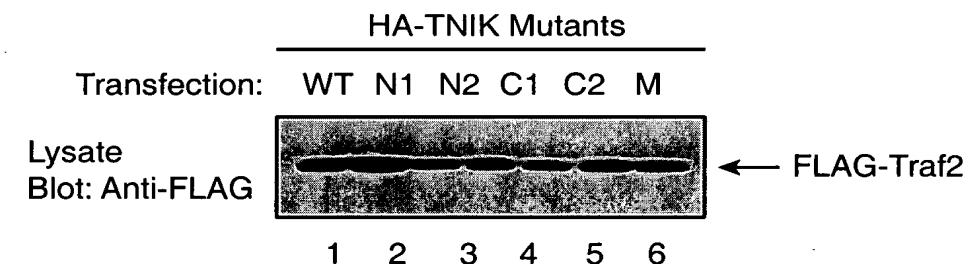


FIG._8C

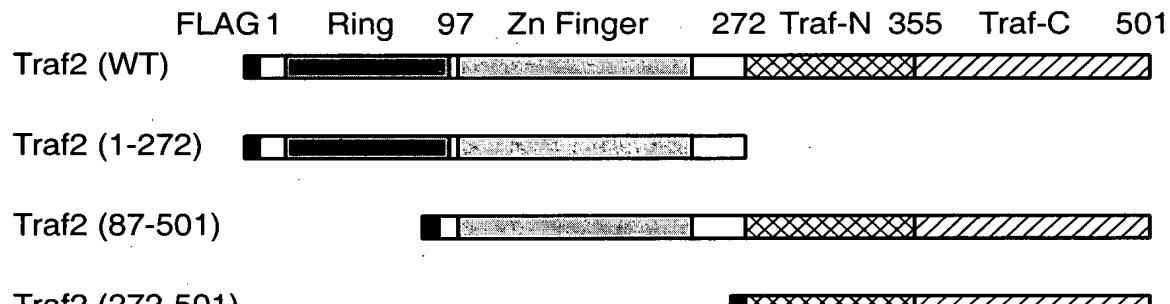
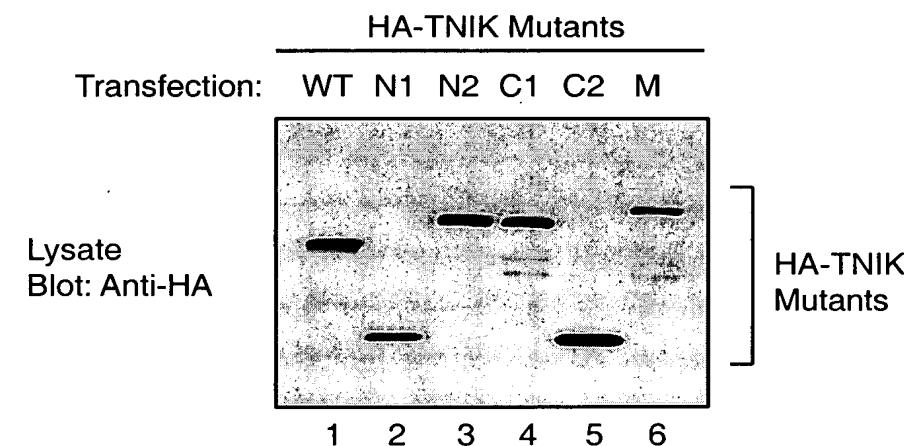
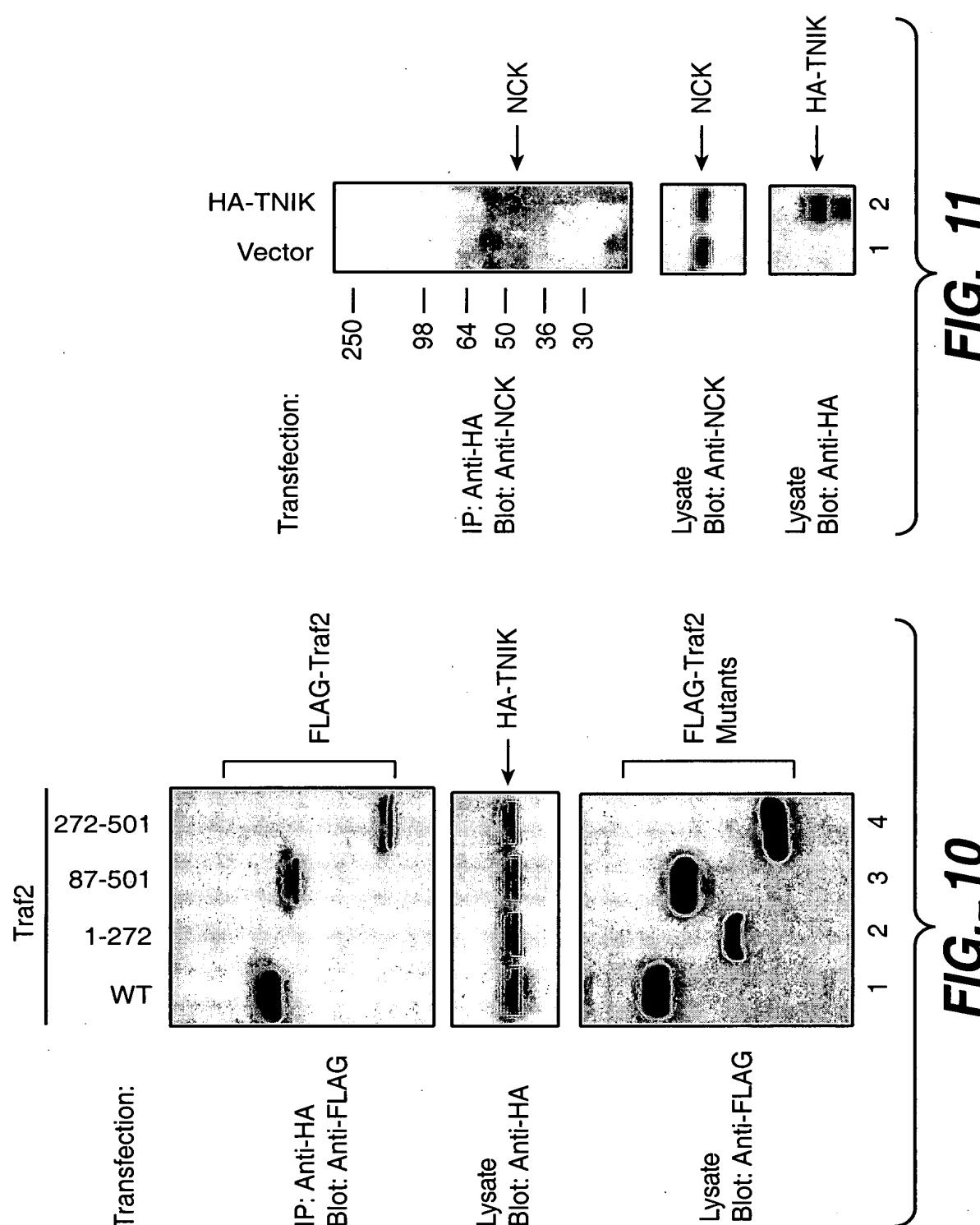
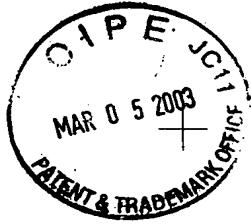


FIG._9





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FIG._12

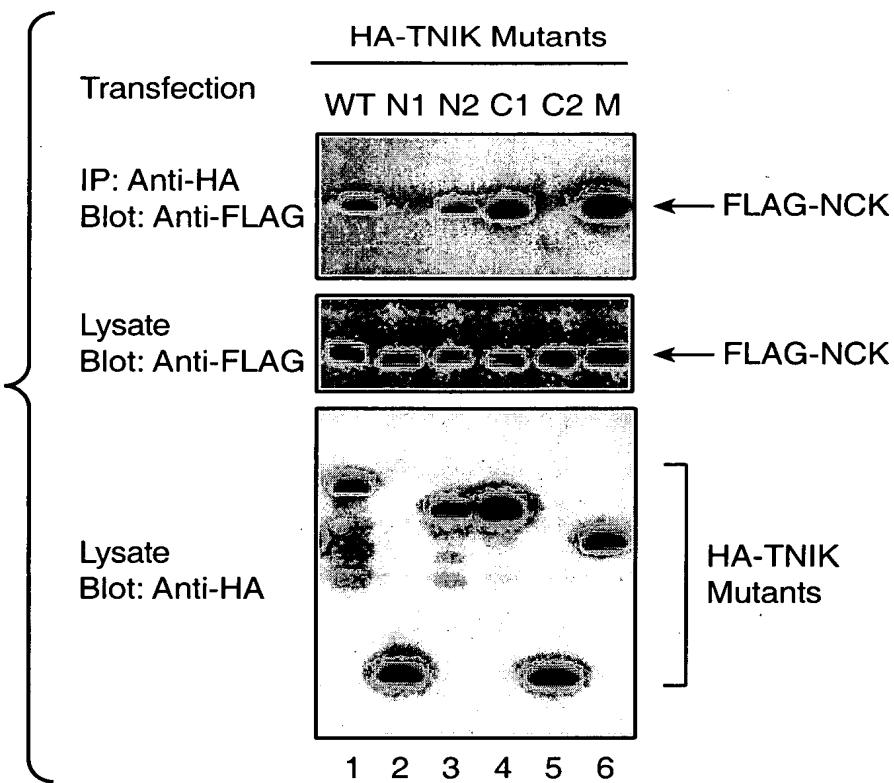
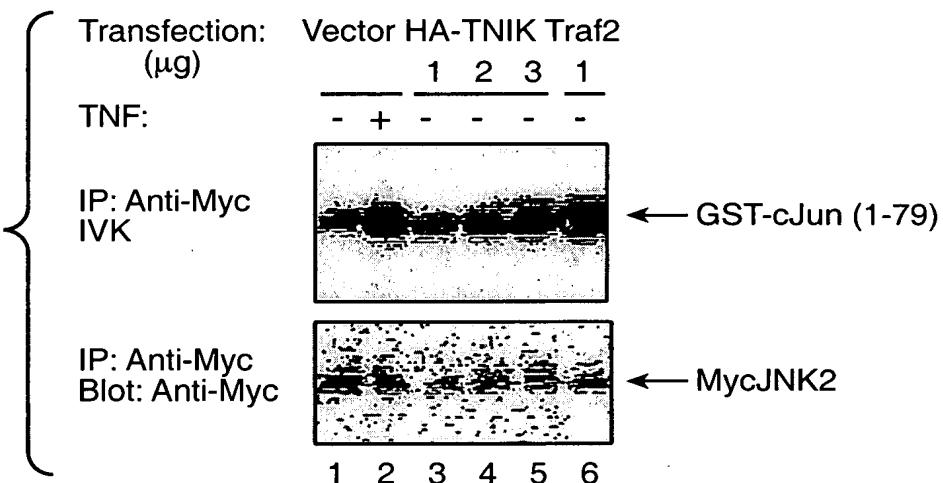


FIG. 13





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FIG._ 14

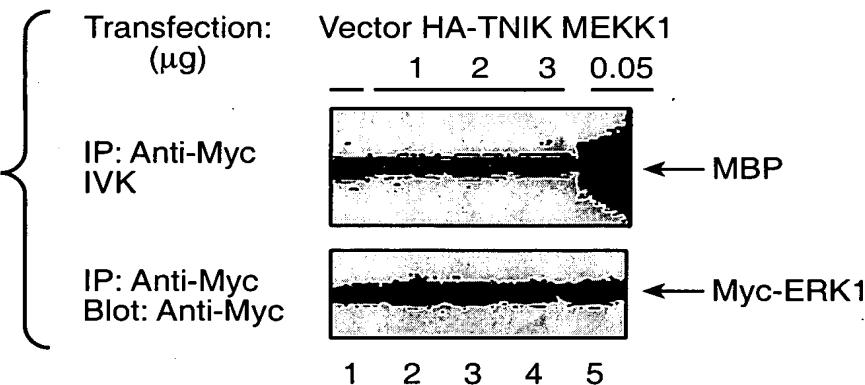


FIG._ 15

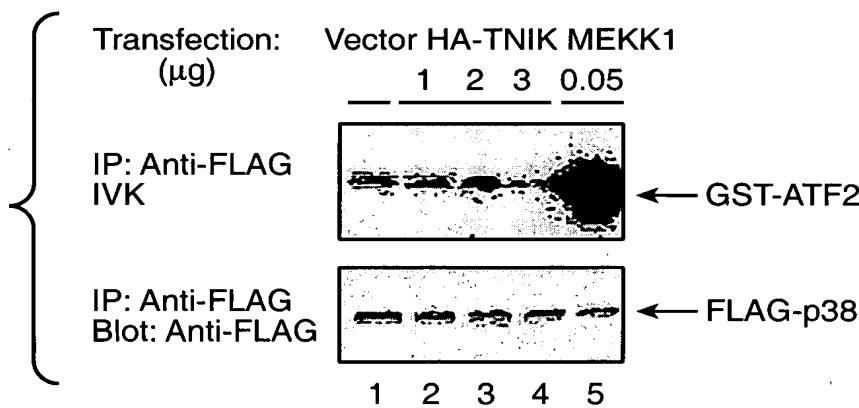
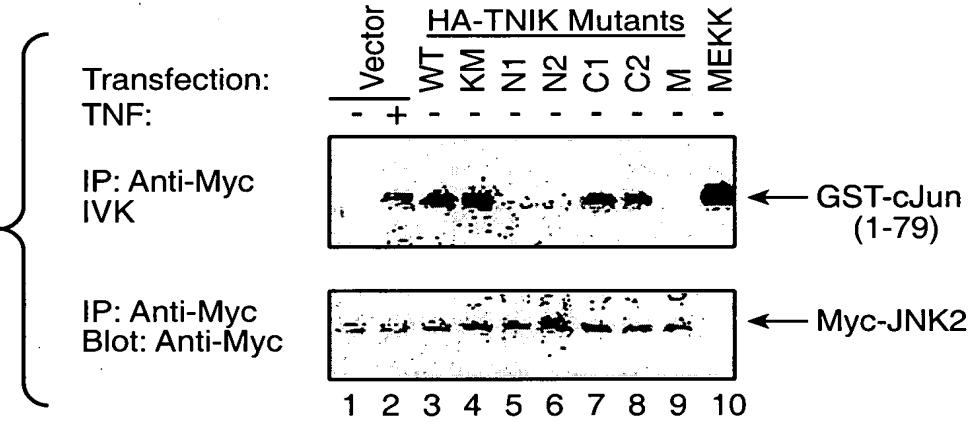


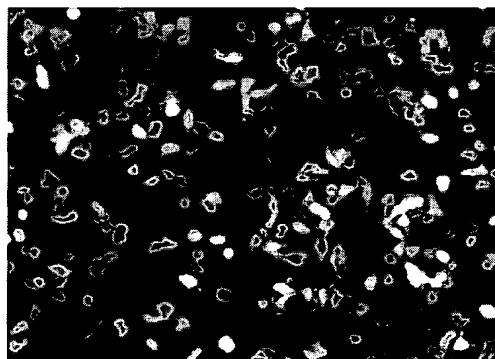
FIG._ 16



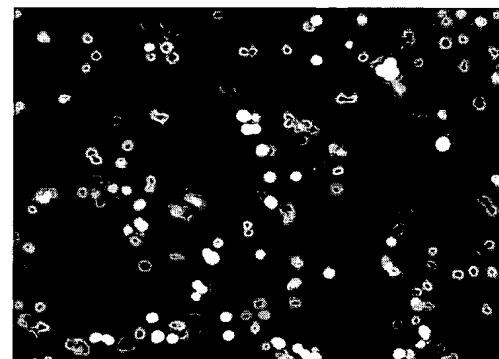


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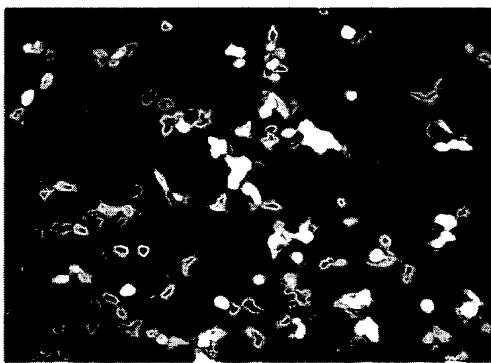
Vector



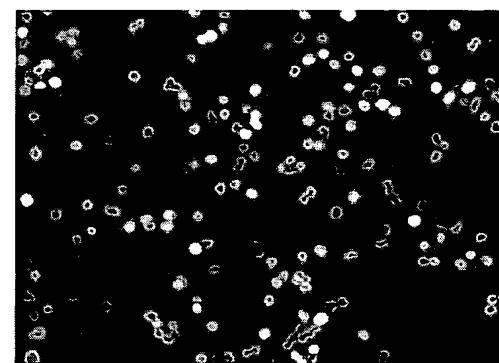
TNIK



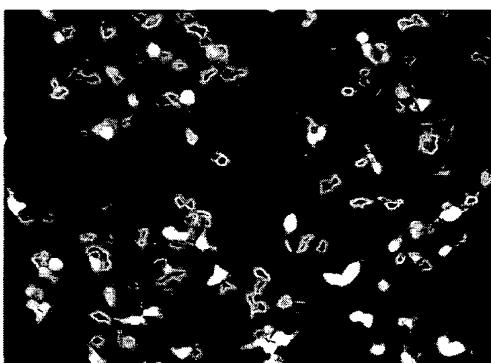
TNIK (KM)



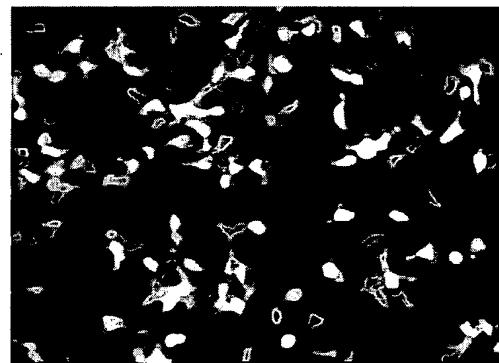
TNIK (N1)



TNIK (C1)

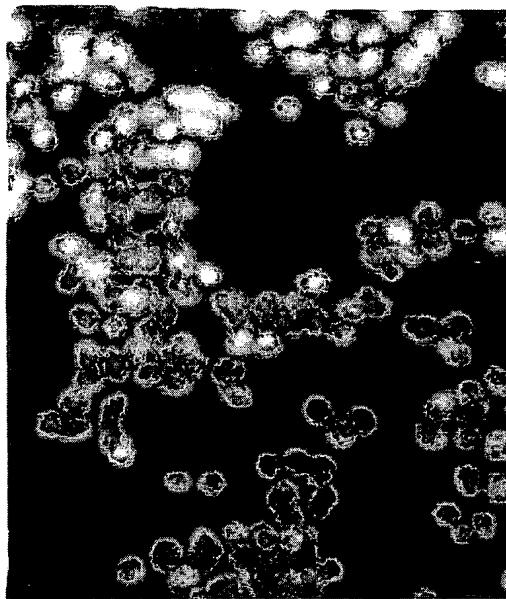


JNK2

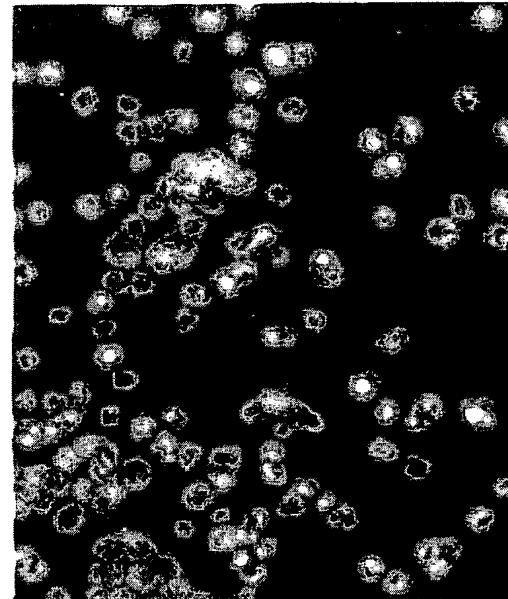
**FIG._17**



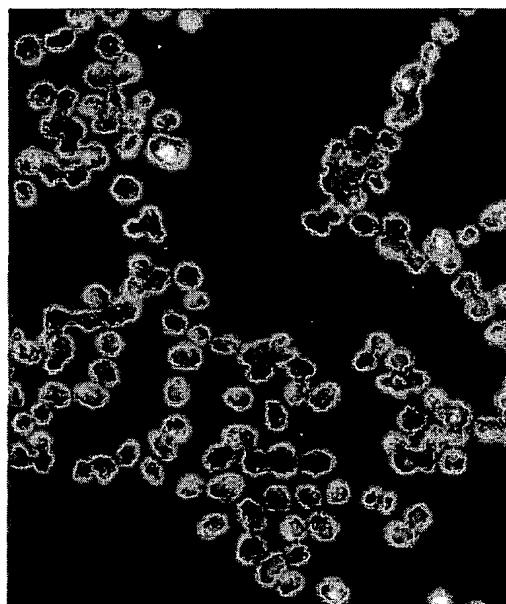
TNIK (WT)



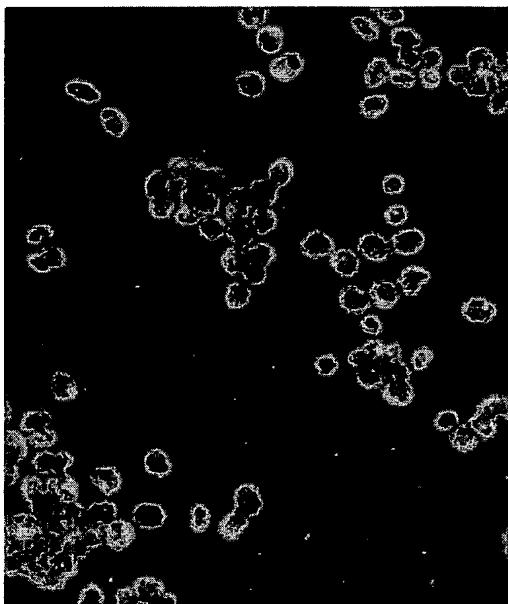
RIP

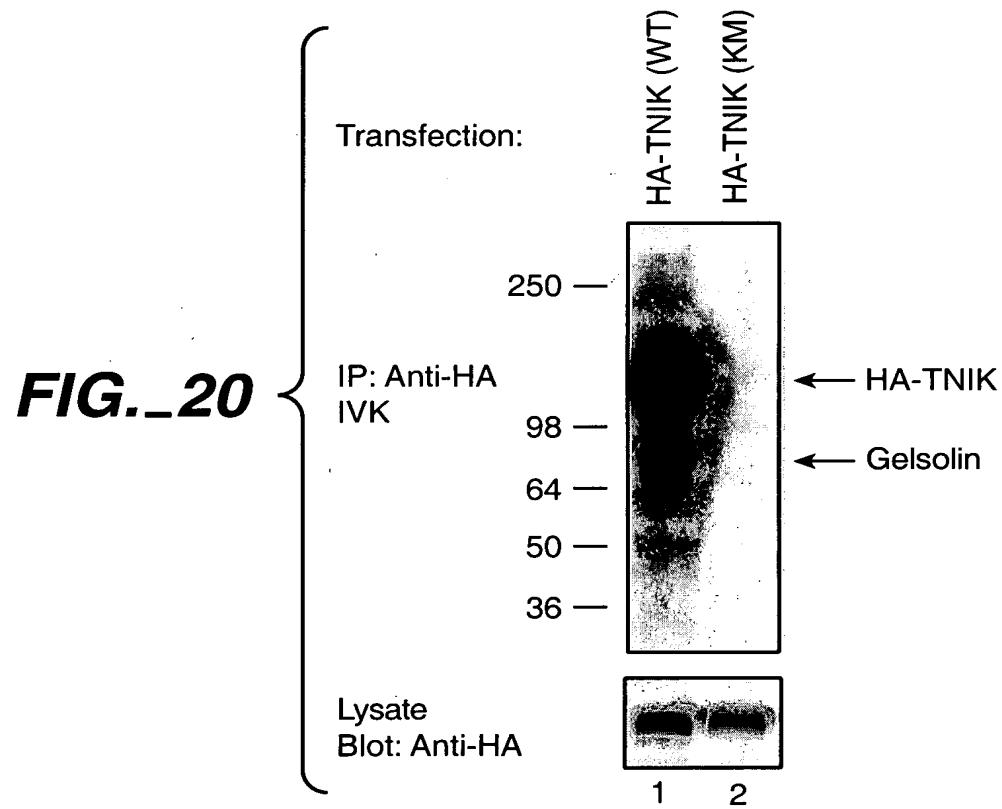
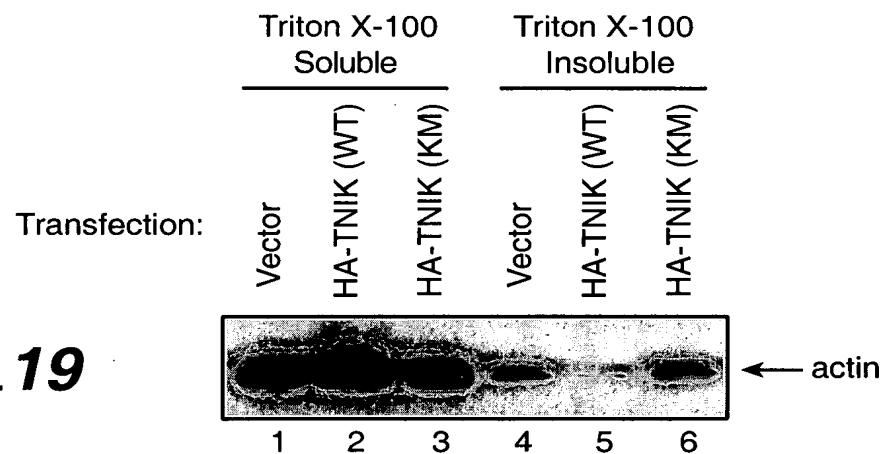


Vector



TNIK (KM)





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FIG. 21A



ACAACTTCTATATCCCCAGCATTAGCCAGAAAGAATTCTCCTGGAAATGGTAGTGCTCTGGGACCCAGACTAGGA
TCTCAACCCATCAGAGCAAGCAACCCCTGATCTCCGGAGAACTTGAGCCATCTGGAGGCCCTGGCAGAGGAC
AGCAGTGGCAGTTCCAGGCTCCAGCTCCAGCACCCTAGCTCCAGCACCCTAGCTCCAGGGCTCCAGCAGGCTGGATCA
CAAGCAGGATCCAGTGAACGCCAACAGAGAGAGGTTCAAGTCAAGAGGATCACCCTGGCTACCTGTGCTTCCCAGATGAG
CCTGGCCAAAGGTGAACCCAGAAATCCAGGGACATTACCCGGGACATTACCCGGGACATTACCCGGGACATTACCCGGGAC
GATGAGGGATCTGACGGCATTAGCCAAAGAAGTAAAGGAGAACTCCGGGATGAGGGATCTGACGGCATTAGCCAAAGAAG
GTGACTGATTACTCCTCCCTCAGTGGAGGTAGAAGTGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG
CATGATGGGACAGTGGCTGTCAAGGCACATACCCAGACTGATAACCAACAGGAGCTCCAGGCAGCAACGAGCAGTAC
AATGTTGGGATGGTGGAGATGGGACTCTGGGCTGGAGGACCTCTGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG
GGAACCTTGTGATGATTAGAGAGACGCTCTGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG
CACATCAACCTCCCTGACCTGGGACTCTGGGCTGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG
TCAACCCATCCCAGGGAG
TTTGTTGGACCCCCAGAGTATACCCAGACGGTCTCCACTGATGAAAGATGAGGAGGAGGAGGAGGAGGAGGAGGAG
CTGTTTACTAGCGAACCTCTTAGGCAAGAACAGGCCAAACTCAATGAAAGCAAGAAAGGATTGGTGGTAAATGTA
AACCCAACCAACATTGGGCCCTCATAGCGACACACCAGAAATCAGAAAATACAGAAAACCGATTGGCCTGATGCTTGG
CTTGTGCAGCTCTGTGGGGTGTAAACCTTCTGGTGGGGACTGAAAATGGCCTGATGCTTGGGGACTGAACTGGCT
CTGTTTACTAGCGAACCTCTTAGGCAAGAACAGGCCAAACTCAATGAAAGCAAGAAAGGATTGGTGGTAAATGTA
AACCCAACCAACATTGGGCCCTCATAGCGACACACCAGAAATCAGAAAATACAGAAAACCGATTGGCCTGATGCTTGG
CAAGGGCAAAGTCTATAATCTGATCAACCGAGGGGATTTCAGCAGATGGATGCTAGAGGGACTGAACTGGCT
GTGACAATTTCAGGAAAGAAGAATAAGCTACGAGTTACTATCTTCTATGGTTAAGGAAACAGAAATACATAAAT
GACCCAGAAGTGGAAAGAACAAAGGCTGGATCACTGGGACTTGGAAACTTGGGACTTGGGACTTGGGACTTGG
AAATATGAAAGGATCAAATTGGTGTGATTGGCCTTAAGGAATGCTGTGGAAATATATGCTTGGGCTCCTAAACCG
TATCATAAATTCAATGGCATTAAAGTCTTGCAGATCTCCAGCACAAAGCCTCTGCTAGTTGATCTCAGGGTAGAA
GAAGGTCAAAGATTAAAGGTATTGGTTCAACACTGGTTCCATGTAATTGATGTTGATCTAGGGATAACTAAGGATGT
TATGATATCATACCATCTCATATTCAAGGGCAATATCAGGGCAATATCAGGGCAATATCAGGGCAATATCAGGG
GGAATGGAAATGCTTGTGCTATGGGGATGAGGGGTGTATGTAACACACTGGTTCCATGTAATTGATGTTGATCTAGGG
GGCTCCAAATGGGGAGAAATGCCAACAGGTCTGGCCTACATTCAATTCACTCCCTCATGCTTATGCTTGGGCT
GCTATTGAGATCCGGTCAGTGGAAACAGGACATTGGATGGAGTATTATGCTAAAGGGTAAAG
TTTCTATGTTGAAGAAAGGAAATGATAAGGTATTGGCATCCGGGATCTGGGAGGAAAGTAGGCCAAGTGGTTCAAG
ACCCCTCAACAGAAATTCATGATGAACTGGTAA

FIG.-21B



FIG. - 22A

TCCCGAGCATTAGCCAGAAAGAAATTCTCCTGGGAATAGGTAGTGGCTCTGGGACCCAGACTGGATCTCAACCCATC
 AGAGCAAGCAAACCCCTGATCTCGGAGAACTGAGCCATCTGGGACCCCTAGCTCCAGCCCCAGCTCCAGGAGGATCC
 TCCTCCAGCTCCAGCACCCCTAGCTCCAGCCCCAGCTCCAGGAGGATCACAGTAAGTCAGAAGGATCACAGGAGGATCC
 AGTGGAAACGCAACAGAGGATCACAGTAAGTCAGGAGGATCACAGTAAGTCAGGAGGATCACAGTAAGTCAGGAGGATCC
 AACACAGAAGAAATCCAGGGACATTACCCGGCCAGCTAGCTACAAAAAGCTATAGATGAGGATCTG
 ACGGCAATTAGCCAAAGGAAACTAAGAGAACTAAGAGAACTCCGGATTGAAGAAACAAACGGCCAAATGAAGAAGGGTGA
 CTGAGGATCTCCAGTGGAGGAGTCAAGAAGGAGTCAAGAAGGAGTCAAGAAGGAGTCAAGAAGGAGTCAAGAAGGAGT
 GTGGGGACCCATGGGCTGGAGACCTCTCATGGGGACAGTGGGAGCAGTGGGAGCTGGGAGCTGGGAGCTGGGAGCTGG
 ATTAGAGAGAGACGGTCTGGAGAGAAGAGGAGTCAAGAAGGAGTCAAGAAGGAGTCAAGAAGGAGTCAAGAAGGAGT
 CCTGACCTGTTGCGAGCAGGCCATTCTCCAGCTGGGAACCCCCGACTGAGGGACTGGGAGCTGGGAGCTGGGAGCTGG
 CAGGGAGATGGACTCTGGGACTGAAATGGCATGGGAGCAGGCCAAAGCCTCCTTCACCCCTGGGAGCTGGGAGCTGG
 AGAGTATACCAGACGGTCTCCACTGATGAAGGATGAAGGAGTCAAGAAGGAGTCAAGAAGGAGTCAAGAAGGAGTCA
 GAACTTCTTAGGCAAGAACGGCCAAGAACGGGAAACTCAATGAAGGAAAGATTGGGAGCTGGGAGCTGGGAGCTGG
 ATTGGGCTCATAGGGACACACCAGAAATCAGAAAATACAAGAAACGGATTCAACTCAGAAATACTTGTGGCAGCT
 CTGGGGTGTAAACCTTCTGGGGGAGCTGGAAAATGGCCTGATGCTTTGGGACCGAAGGTGGGAGCTGGGAGCTGG
 TATAATCTGATCAACGGAGGGGAGATTTCAGCAGATGGATGCTAGAGGGACTGAATGTCCTTGTGACAAATTCA
 GGAAAGAAGAAATAAGCTACGGAGTTACTATCTTCATGGTTAAGAAAAGAAATACTACATAATGACCCAGAAAGTA
 GAAAAGAAAACAAGGGCTGGGATCACTGGGGGAGCTGGGAGCTGGTGTATACATTATAAGGTGTTAAATGAAAGG
 ATCAAATTTGGTGTGGCCTTAAGGAATGCTGTGGAAATATGCTTGGGCTCCTAAACCGTATCATAAATTCA
 ATGGCATTAAAGTCTTTGCAAGATCTCCAGGCCAACGGCCTCTGCTAGTTGATCTCAGGGTAGAAGGAAGGTCA
 TTAAGGTTATTTGGTTCAACACACTGGTTCCATGTAATTGATGTTGATTCAAGGAAACTCTTATGATATCTAC
 ATACCATCTCATATTCAAGGCAATATCACTCCTCATGGCTTAAACACACTTAAGGATGTGGTGTGGCTCAATG
 CTTGTTGCTTATGAGGATGAGGGGGTGTATGTAAGGATGGGGCTTCAAGGAAACTCTTATGATGAAATGGA
 GGAGAAATGCCAACGGTCTGGCCTACATTCAATTCAGATAATGGGCTGGGGAGAAAGCTATTGAGATC
 CGGTCAAGTGGAAACAGGACATTGGAGTGGAGTATGCAATAAGGGAGCTCAAAGGTTAAAGTTCTATGTGAA
 AGAAATGATAAGGTTATTTGGCATCCGGGATCTGGGAGTGGAGTGGAGTGGAGTGGAGTGGAGTGGAGTGG
 AATTCCATGTAAGTGGTAA

FIG.-22B

FIG. 23A



FIG. - 23B



ATGGCGGAGCCACTCCCCGGCTCGAAGGCCATTGGATGAAAGATGATCTCGGGCTCTGAGGGACCCCTGCGAGGGATCTT
 GAATTGGGTGGAACCTGGTTGGAAATGGAAACATACGGGCAAGTTATAAGGGTCTGGTCATGTCACAAACGGGCGAGCCTT
 GCAGGCCATCAAGGTATGGATGTCACAGGGATGAAAGAGGAACATCAACAGAAATTAAACATGTTGAAGAAA
 TATTCTCATCACCGGAATTGGCTACATGGCTACATGGCTTATCAAAAGAACCCACAGGCATGGATGACCAA
 CTTGGTGGTGGATGGAGTTTGTGGATGTCACGGACCTGATCAAGAACACAAAGGTAACACGGCATGGATGACCAA
 AAAGAGGAGGTGGATGGCATACATCTGCAGGGAAATCATCTGCAGGGAAATCTTCATGGGAACCTGGCACCAGCATAAAGGTGATT
 CATCCGAGATTTAAGGGCAAATGTCCTGGCTGACTGAAAGTTAAACTAGGGACTTTGGAGTCAGTGGATGGCACCAGAAGGTTATT
 GCTCAGCTGATCGAACAGTGGCAGGGAAATACTTCATGGGAACCTCCCTACTGGATGGCACCAGAAGGTTATT
 GCCTGTGATGAAACCCAGATGCCACATATGATTCAAGAGTGACTTTGGGTCTTTGGGTATCACCGCCATTGAA
 ATGGCAGAAGGTGCTCCCCCTCTGTGACATGCACTGGAGGCTCTCCTCATCCCCGGAAATCCAGGCC
 CCTCGGCTGAAGTCTAAGAAGTGGTCAAAAATTCCAGTCATTGAGAGGTGGTAAAGAAATCACAGC
 CAGCCGACCGAAACAGAAACAAATTGATGAAAGCATCCATTATAGAGACCAACCTAATGAGGCACAGGTCAGGTGGCATT
 CAACTCAAGGGACCATATTGATAGAACAAAGAAGGAGGGAAAAGGATGAGACAGAGTATGAGTACAGTGG
 AGTGGAGAAGGAGGGAGGAAATGACTCAGGAGGCCAGCTCCATCCTGAATCTGGCCAGGGAGTGGCAGGCTG
 CGGAGGGACTTTCTGAGGCTGGCTGAGGCTGGCAACAAAGGAGGGTCTGAGGGCCTACGGAGGGCAGCTGGAGG
 CAGCAGGGGAGAATGAGGAGCACAGGGCAGCTGGCCAGGGCAGCTGGAGGAGGAGGAGGAGGAGGAGG
 CAGAGGGGGGGCTGGAGGAGCAACAAAGGCCAGGGAGGGAGGAGGAGGAGGAGGAGGAGGAGGAGG
 CACTATGAGGGAGGAGGAGACAGTTAGAGATGCGGCCAGGGAGGAGGAGGAGGAGGAGGAGGAGG
 TTAGAGGGAGGAGGAGGAGACAGTTAGAGATCTGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGG
 AAGCCAAACAAATTGGAAGAACAGAGACAAGGAGAAAGGAGAAAGGAGGAGGAGGAGGAGGAGGAG
 GTTCCCTTCAAGCATCGGGGGCAGGGAGCAGGGCCTGTGGAGGAAGGCCACTGTACAGCTAAAGGAAGACTTA
 AGTCCTAGTGTGAGAACGCCAGGGAGGTAGAACGGTCAAGGGCAAGGCTCAACCGGAAAGGTTCCCTGCC
 ATGCCCTCACAGGTGCCAACAGGATATCTGACCCCCAACCTGGAGGAGTCCCACAGTCAGATCCCACATCTGGTA
 GTTCAGGCTGCTGAACACCCCCATGCTCAGACCCAGTCAGGAGGAGGAGGAGGAGGAGGAGGAGG
 CAGGGACCTGCCCTGGACGGCCAGGTGACGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGG
 CTGAAACGTGACCTCCACCGCGTGGAGATGCCACGGAGGAGTCCCACAGTCAGATCCCACATCTGG
 ACTCGCATTGAAAGGTTAGCAGACGGAGAACATCCACCAAGGGCCTCAAAAG
 ACAACTTCTATCCCCAGCATTAGCCAGAAAGAATTCTCTGGGAATGGTAGTGTGCTGGGACCCAGACTAGGA

FIG.-24A

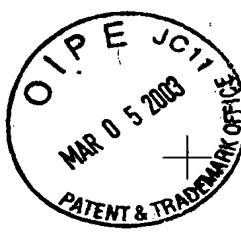


FIG. - 24B

1

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FIG. - 25A



GTTCGAGCCAACAGTAAAGTCAGAAGGGATCACCTGCTTCCCCATGAGCCTGCCAAGGGTGAACCGAGAAATC
AGGGACATTACCCGGCCAGTCGACCCAGCTAGCTACAAAAAGCTATAAGTGAAGGGATCTGACGGCATTAGCCAAA
GAACTTAAGGAGAACTCCGGATITGAAGAAACAAACCGCCCAATGAAAGAAGGTGACTGATTACTCCTCCTCCAGTGAG
GAGTCAGAAAGTAGCGAGGGAAAGATGGAGAGGGAGACCCATGATGGGACAGTGGGACAGTGGGAAATGGGGCATGGCAG
ATACCCAGACTGATACCAACAGGAGCTCCAGGCAGCAACGGAGCTAACAGGAGCTAACAGGAGCTAACAGGAGCTAACAG
CTGGAGACCTCTCATGGGACAGTTCAAGAGAACCTTGTATGATTAGAGAGACGTCT
GGAGAGAAAGCGATCTGGCCACAGTGACAGCAATGGCTTGGCTGGCCACATCAACCTCCCTGACCTGGTGCGAG
CAGAGCCATTTCTCCAGCTGGAACCCCGACTGAGGGACTGGGGACTGGCGCTCCTCAACCCATTCCAGGAGATGGACTCT
GGGACTGAAATATGGCATGGGAGCACCACAGCCTTCACCCCTTGTGGACCCCAGAGTACCAAGAGC
TCTCCCACTGATGAAGAGTGAAGGGATGAGGAATCATCAGCCGAGCTCTGTACTAGCGAAACTTCTAGGCCAA
GAACAGGCCAAACTCAATGAAGCAAGAAAGATTGGTAAATGTAACCCAAACCPAACATTGGCCTCATAGC
GACACACCAGAAATCAGAAAATACAAGAAACGATTCAACTCAGAAATTAACCTTGTGCAGCTCTGGGGTGTAAAC
CTTCTGGTGGGACTGAAAATGGCCCTGATGCTTTGGACCGAAAGTGGCAAGGGAAAGTGGGACTGAAATGTCCCTGTGACAA
CGGAGGGGATTTCAGCAGATGGATGTGCTAGAGGGACTGAAATGGTACATTTCAGGAAGGAATAAG
CTACGAGTTACTATCTTCTATGGTTAAGAACAGAATACATACATAATGACCCAGAAGTAGAAAGAACAAAGGC
TGGATCACTGTTGGGACTTGGAAAGGCTGTATACTATAAGTGTGTTAATATGAAAGGATCAATTTCATGGCATTAAAGGTATT
ATTGCCCTAAAGAATGCTGTGAAATATGCTTGGCTCCTAACCGTATCTACAGGTTAGAAGGTCAAGGTTAATTTGGTG
TTTGCGAGATCTCCAGCACAGCCCTCTGGCTAGTTGATCTCACGGTAGAAGGTCAAGGTTAATTTGGTG
GGTTCACACACTGGTTCCCATGTAATTGATGTTGATCTGCTTGGCTAAACAGATGGAAATGGAAATGCTTGGCTATGAG
CAGGGCAATATCACTCCCTCATGCTTGGCTAAACACCTATGGCCGGATAACTAAGGGATGTGGCTCCAAATGGGAGAAATGCCCAAG
GATGAGGGGGTGTATGTAACACCTATGGCCGGATAACTAAGGGATGTGGCTCCAAATGGGAGAAATGGGAAAGCTATTGAGATCCGGGTCAAGTGGAA
TCTGTGGCCTACATTCAATTCAATCAGATAATGGGCTGGGGAGAAAGCTATTGAGATCCGGGTCAAGTGGAA
GGACATTGAGTGGAGTATTATGCATAAGCGAGCTCAAGGTTAAAGGTTCTATGTGAAAGAAATGATAAGGTA
TTTTTGCATCCGTGGGATCTGGAGGAAGTAGCCAAAGTGTGTTTCATGACCCCTAACAGAAATTCATGATGAAC
TGGTAA

FIG._25B



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FIG. 26A

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FIG. 26B



FIG. 27A



FIG. 27B

1



ATGGCGAGGCCACTCCCCGGCTCGAAGGCTCTGGATGAAATAGATCTCTGGCTCTGGACCCCTGCAGGGATCTT
 GAATTGGTGAACCTGGAAATGGAACATACGGGCTTATGGATGTCACAGGGATGAAGGAGAAATCAAACAGAAATTAAACATGTTGAAGAAA
 GCAGGCCATCAAGGTTATGGATGTCACAGGGATGAAGGAGAAATCAAACAGAAATTAAACATGTTGAAGAAA
 TATTCTCATCACCCGAATATGCTACATACTATGGTCTTATCAAAAAGAACCCACAGGATGGATGACCAA
 CTTGGTTGGATGGAGTTGCTGGCTCTGGCTCACCGACCTGATCAAGAACACAAAGGTAAACACGTTG
 AAAGAGGAGGATGGATTGCATACATCTGGCAGGGAAATCTTACGGGGCTTGAGTCACCTGACCCGATAAAGTGTAT
 CATCGAGGATATAAGGGCAAATGTCCTGGTGAACGAAAATGAGAAGTTAAACTAGTGGACCTTGGAGTCAGT
 GCTCAGCTTGTATCGAACAGTGGCAGGGAAATACTTCATGGAAACTCCCTACTGGATGGCACCAGAACGTTATT
 GCCTGTGATGAAAACCCAGATGCCACATATGATTCAAGAGTGAACCTGTTGGCTTTGGGTATCACCGCCATTGAA
 ATGGCGAGGGTGGCTCCCCCTCTGTGACATGCACTGGCACCCATGAGAGCTCTCTCCTCATCCCCCGGAATCCAGCG
 CCTGGCTGAAGTCTAAGAAGTGGTCAAAAAAAATTCCAGTCATTATTGAGAGCTGTGCTGGTAAAGAAATCACAGC
 CAGCGACCAGCAACAGAACATTGATGAAGCATCCATTATACGAGACCAACCTAATGAGGACAGGGACAGGGTCCGCATT
 CAACTCAAGGACCATATTGATAAGAACAAAGAACAGGAGGAGAAAAGATGAGACAGAGTATGAGTACAGTGGAA
 AGTAGGGAAGAGGAGGAATGAACTCAGGAGAGGCCAGCTCCATCCTGAAATCTGCCAGGGAGTGGCAGCGCTG
 CGGAGGGACTTTCTGGGCTGGCTGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG
 CAGCAGCGGGGCTGGAG
 CACTATGAGGAGGAGATGCCAGATGG
 TTGGAAAGAACAGAGAACAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG
 CATCAGCGGGAGGGAG
 AAGCAGCATGGGCAAGGAGATCCCACATCTGGTAGCTGTAATCCAGGGACCTGCCTGACCGCCCTCCAG
 TCAGTGCACGGCAGGCCAACAGGGCCTCTGGGTTTCAGGGGGCTCTGAACGTGACCCCTCCACCTGGCATTGAAAGTTGACCGAAGC
 ATGCCACGCCCAGATCCCACCTGGAAAATCCTCCACTGGCATTGAAAGAACACTTCTATATCCCCAGCATTAGCC
 TCTGGTTACGACAGGAAGAACATCCACCAAGGAGACATTCCACCAAGGAGGAGGAGGAGGAGGAGGAGGAGGAG
 AGAAAGAATTCTGGGAATGGTAGTGGCTGGGACCCAGACTAGGATCTCAACCCATCAGAGCAAGCAACCCCT
 GATCTCCGGAGAACTGAGGCCATCTGGAGAGGCCATCTGGAGAGGCCAGCTGGGAGGAGGACAGGGCAGTGGGATCACAGGAC
 ACCCCTAGCTCCAGGCCAGGCTCCCAAGGAGGCTCCAGGCCACAGGAGGATCCAGGAC

FIG._28A



GTTCGAGCCAACAGTAAGTCAGAAGGGATCACCTGCTGGCTTCCCATGAGGCCAAGGTGAAACCAGAAGAAATCC
AGGCACATTACCCGGCCAGTCACCGCTGATCTGACGGCATTAGCCAAGAAACTAAGAGAAACTCCGGATTGAA
GAGGAAGAGATGGAGAGGGACCCATGATGGGACAGTGGCTGTAGGGACATACCCAGACTGATAACCAACAGGA
GCTCCAGGGCAGCAACGGGAGCTACATGTGGGAATGGTGGGACCTCTCATGGGGACAGT
TCAGGGGCACTATTCAAGAGAAGGGAAACCTGTGATTAGAGAGCCTGAGAGAAGGAAGGGATCTGGCCAC
AGTGACAGGCAATGGCTTGGCCACATCACCTCCCTGACCTGGTGCAGCAGGGCATTCCAGCTGGAAAC
CCGACTGAGGGACTGGGGCGCTCTCAACCCATTCCAGGAGTACCCAGGATGGACTCTGGACTGAAATGGCATGGGAGC
AGCACCAAGCCTCCCTCACCCCTTCAAGGGGACCCAGAGTACCCAGGATGGACTCTGGTCCCAACTGAAAGATGAAGAG
GATGAGGAATCATCAGCCGAGCTCTGGTACTAGCGAACTCTTAGGCAAGAATGGGAAACAGGCCAAACTCAATGAAGCA
AGAAAGATTTCGGGTAAATGTAACCCAAACATTGGCCTCATAGGACACACAGAAATCAGAAAATAC
AAGAAACGATTCAACTCAGAAATACTTTGTGAGCTCTGGGGGTAAACCTTCTGGGGGACTGAAAATGGC
CTGATGCTTTGGACCGGAAGTGGCAAGGCAAAGTGGCTCTGGACAATTTCAGGAAAGAAATAAGCTACGAGTTACTATCTTCATGG
TTAAGAAAACAGAATACTACATAATGACCCAGGAAGTAGAAAAGAAACAGGCTGGATCACTGTTGGGACTTGGAA
GGCTGTATAACATTAAAGGTGTTAAATATGAAAGGATCAAAATTGGCATTAAAGTCTTGGCAGATCTCCAGCACAGGC
ATATATGCTGGCTCCTAAACCGTATCATAAATTCAATGGCATTAAAGGTTCAAGGTTAGAAAGGATTAAGGTTATTGGTTCAACAC
CTGCTAGTTGATCTCAGGGTAAAGGATCTCAGGAAACACTCTTGTGATATCTACATACCATCTCATATTCAAGGGCAATATC
ATTGATCATCTGGCTAAACAGATGGAATGGAAATGCTTGGCTATGAGGATGAGGGGTGTATGTAACAC
TATGGCGGATAACTAAGGATGTTGGCTCCATGGGAGAATGCCACAGTCTGTGGCCTACATTCAATTG
CAGATAATGGCTGGGGCGAAAGCTATTGAGATCCGGTCAGTGGAAACAGGACATTGGATGGAGTATTG
CATAGCGAGCTCAAGGTAAAGTTCTATGTAAGGAAATGATAAGGTATTGCTTGCATCCGGATCTGGA
GGAAGTAGCCAAAGTGTTTCATGACCCCTAACAGAAATTCCATGAAACTGGTAA

FIG.-28B



1 MASDSPARSILDEIDL SALRD PAGI FEL VEL VNG NG TY QQVY KGRHV KTGQLAAIKVMDVTG
61 DEEE EIKQE IMLKKY SHRNIA TTYG AFI KKNPPGMDDQLWLVM EFCGAGS VTDL I KNT
121 KGNTLKEEWIA YICREI LRLGLS HLLHQH KV IHRDIK GQNV LLEN TMAEVKL VD FGVSAQLDR
181 TVGRRNTFIGTPYUWMAPEVIACDENPDATYDFKSDLWLSL GITAI EMAEGAPPLCDMHPMR
241 ALFLIPRNPAPRLKSKWSKKFQSFIESCLVKNHSQR PATEQLMKHPFIRDQPNERQVR
301 QLKDHIDRTKKRGEIDETEYEYSGSEEEENDSGEPSSILNLPGE STLRRDFLRLQLA
361 NKERSEALR RQQL EQQ RENE EH KQLLAERQKRIEEQKEQRRL EEEQRRREKE L RQQE
421 REQR RHYEEQMR REEERRRAEHEQY KRKQ LEEQRQAERL LQRQLKQERDYL VLSLQHQRQE
481 QRPVEKKPLHYKEGMS PSEKPAWAKEE VERSRLN RQSSPAMPHK VANRISDPNLP PRSE
541 SFSISGVQPARTPPMLRPVDPDQIPHLVAVKSQGPALTASQSVHEQOPTKG LSGFQE ALNVT
601 SHR VEM PRQNSDPTSEN PPLPTRIEKFD RSSWL RQEE DIPP KVPQRTT SISPA LARKN SP
661 GNGS ALG PRLGSQPIRASNPDLIRRTEP ILESPLI QRTSSGSSSSSTPSSQ PSSQ GSGQ P
721 SQAGSSERTRV RANSKSEGSPVLPHEPAKVKPEESR DITRPSR PASYKKA IDEDL TALAK
781 ELRELRIEETNRP MKKVTDYSSSEESSEESSEEEEDGESEETHDGTVA VSDIPR LPTGAP
841 GSNEQYQNVGMVQGTHGLETS HADSFSGSISREGTLMIRETSGEKKRSGHS DSNGFA GHINL
901 PDLVQQSHSPAGTPTEGLGRV STHSQEMDSGTEYGMGSSTKASFTPFVDP RVYQTSPTDE
961 DEEDEESSAA ALFTSELLRQEQAKUNNEAR KISV VVNPNPTNIRPHISDTPEIRK YKKRFNSE
1021 ILCAALWGVNLLVGTENG LMLLD RSGQGK VYNL INRRRFQ QMDVLEG LNV LVTISG KQNK
1081 LRVY YLSWL RNRILHNDPEVEKKQGWITVG DLEGCIHYK VVKYERIKFLVIA LKNAVEI Y
1141 AWAPKPYH KFMARKSFADLQH KPLLV DLTVEEGQRL KVIFGSHTGFHV IDV DSGNSYDIY
1201 IPSHIQGNITPHAI VILPKTDGMEML VCYEDEGV VNTYGRITKD VV LQW GEMPT SVAYI
1261 HSNQIMGWGEKAIEI RSVETG HLDGV FMHK R A QRLKFL CERNDK VFFAS VRS GGS SQVFF
1321 MTLNRNSMMNWZ

FIG.-29



1 MASDSPARSLEIDLSALRDPAGIFELVELVGNNGTYGQVYKGRHVKTGQI LAIKVMDVTG
 61 DEEEEIKQEINM LKKYSHHRNIA TTYGAF I KKNPPGMDDQLWL VMEFCGAGSVTDLIKNT
 121 KGNTLKEEWIA YICREJLRLGSLH LHQHKV I HRD I KGQNVLLTENAEVKL VDFGVSAQLDR
 181 TVGRRNTFIGTPYMAPEVIA CDENPDA TYDFKSDLWSLGI TAIEMAEGAPPCLCDMHPMR
 241 ALFLIPRNPAPRLLKSKKKFQSFIESCLVKNHSQR PATEQLMKHPPFIRDQPNERQVR I
 301 QLKDHIDRTKKRGEKDETEYSGSEEEEENDSGEPSSILNLPGESTLRRDFLRLQLA
 361 NKERSEALRRQQLEQQQRENEEHKRQLLAERQKRRIEEQKEQRRRLEEQQRREKE LRKQQE
 421 REQRRHYEEQMRREEERRRAEHEQEYI RRQLEEEQRQLEI LQQQLLHEQALLLEYKRKQL
 481 EEQRQAERLQRLQKQERDYLVLSSLQHQRQEQRPV EKKPLHYKEGMSPSEKPAWAKEI PHL
 541 VAVKSQGPALTASQS VHEQOPTKG LSGFQEALNVNTSHRVEMPRQNSDPTSENPLPTRIEK
 601 FDRSSWLRQEE DIPPKV P QRTT SISPALARKNSPGNGSALGPRLGSQPIRASNPDLRRTE
 661 PILESPLQRTSSGSSSSSTPSSQPSSQGGSQAGSSERTRVTRANSKSEGS PVL PHE
 721 PAKVKPEESRDTIRPSR PASYKAIDEDLTALAKELRELRIEETNRPKKVTDYSSSEE
 781 SESSEEEEDGESETHDGTVA VSDI P RLIPTGAPGSNEQYINVGMVGTHGLETSHADS FSG
 841 SISREGTLMIRETSGEKKRSGHSDSNGFAGHINLPDLVQ QSHSPAGTPTEGLGRVSTHSQ
 901 EMDSGTEYGMGSSTKASFTPFVDPRVYQTSPTDEDEEDEESSAAALFTSELLRQEQA KLN
 961 EARKISVNNVNPTNIRPHSDTPEIRKYKKR FENSEILCAALWGVNLLVGTENGMLLDRSG
 1021 QGKVYVNLLINRRRFQQMDVLEG LNLVLTISGKKNKL R VYYL SWL RNR I LHN DPEVEKKQGW
 1081 ITVGDILEGGIHYKVVKYERIKFLVIA LKNAV EYI YAWAPKPYHK FMAFKSFADLQHKPLLV
 1141 DLTVEEGQRLKVI FGSHTGFHVIDVDGSNSYDIIYIPSHIQGNITPHAI VILPKTDGMEML
 1201 VCYEDEGVYVNNTYGRITKDVVLQWGEMPTSVAYIHSNQIMGWGEKAIEIRS VETGHL DG V
 1261 FMHKRAQRLKFLCERNDKVFFASVRSGGSSQVFFM TLNRNSMMNWZ

FIG.-30



1 MASDSPARSLDIEDLSDLRDPAGIFELVELVGNGTYGQVYKGRHVKTGQLAAIKVMDVTG
61 DEEEIKQEINMLKKYSHRNIAITYGAFIKKNPPGMDDQLWLVMEEFCGAGSVTDLIKNT
121 KGNTLKEEWIAVLCREILRGLSHLHQHKVIHRDIKGQNVLLTENAEVKLVDFFGVSAQLDR
181 TVGRRNTFIGTPYMAPEVIACDENPDAFYDFKSDLWSSLGITAIEMAEGAPPLCDMHPMR
241 ALFLIPRNPAPRLKSKWSKKFQSFIESTCLVKNHSQRPATEQLMKHPFIRDQPNERQVR
301 QLKDHIDRTKRRGEKDETEYEYSGSEEEDNSGEPPSILNLPGESTLRRDFLRLQLA
361 NKERSEALRROQLEQQRENEEHKRQLLAERQKRIEEQKEQRRLEEQQRREKELRKQQE
421 REQRRHYEEQMRREEERRRAEHEQEYIRRQLEEEQRQLEILQQQLLHEQALLLEYKRKQL
481 EEQRQAERLQRQLQERDYLVSLOHQHQEQRPVKEKKPLHYKEGMSPSEKPAWAKEVEER
541 SRLNRQSSPAMPHKVANRISDPNLPPRSESFISGVQPARTPPMLRPVDPQIPHLVAVKS
601 QGPALTASQSVHEQPTKGLSGFQEALNVTSHRVEMPRQNSDPTSENPLPTRYIEKFDRSS
661 WLRQEDIIPPKVPRQRTTSISPALARNSPGNGSALGPRLGSQPIRASNPDLRPTEPILES
721 PLQRTSSGSSSSSTPSSQPGSQAGSSERTVRANSKSEGSPVLPHEPAKVK
781 PEESRDITRPSRSPADLTALAKELRELRIETNRPKKVTDYSSSESESEEEEDGES
841 EHDGTVAVSDIPRLIPTGAPGSNEQYNVGMVGTGLETSHADSFGSISREGTLMIRET
901 SGEKKRSGHSDSNGFAGHINLPDLVQQSHSPAGTPTEGLGRVSTHSQEMDSGTEYGMGSS
961 TKASFTPFDPRVYQTSPDDEEDEESSAAAALFTSELLRQEQAQLNEARKISVWNVNPT
1021 NIRPHSDTPEIRKYKRFNSEILCAALWGVNLLVGTENGIMLLDRSGQGKVYNLINRRF
1081 QQMDVLEGLNVLVTISGKKNKLRVYYLSWLRNRIILHNDPEVEKKQGWITVGLEGCIHYK
1141 VVKYERIKFLVIALKNAVEIYAWAPKPYHKFMFKSFADLQHKPLLVDLTVEEGQRLKVI
1201 FGSHTGEHVIDVDSGNSYDIYIPSHIQGNITPHAIILPKTDGMEMLVCYDEGVYVNTY
1261 GRITKDVVILQWGEEMPTSVAYIHSNQIMGWGEKAIEIRSVEETGHIIDGVFMHKRAQRLKFLC
1321 ERNDKVFASVRSGGSSSQVFFMTLNRRNSMMNWZ

FIG.-31



1 MASDSPARSLEIDLSALRDPAGIFELVELVGNNGTYGQVYKGRHVKTGQLAAIKVMDVTG
61 DEEEEIKQEINMLKKYSHRNIAITYGAFIKKNPPGMDDQLWLWMEFCGAGSVTDLIKNT
121 KGNTLKEEWIAIYCIREILRGLSHLHQHKVIRDIKGQNVLLTENAEVKLVDFGVSAQLDR
181 TVGRRNTFIGTPYMAPEVIACDENPDATYDFKSDLWSLGITAIEMAEGAPPLCDMHPMR
241 ALFLIPRNPAPRLKSKWSSKKFQSFIIESCLVKNHSQRPATEQLMKHPFIRDQPNERQVR
301 QLKDHIDRTKKRGEKDETEYEYSGSEEEDSGEPESSILNLPGESTLRRDFFLRLQLA
361 NKERSEALRRQQLEQQQRENEEHKRQLLAERQKRRIEEQKEQRRRLEEQQRREKELRKQQE
421 REQRRHYEEQMRREEERRRAEHEQEYKRKQLEEQRQAERLQRQLKQERDYLVLQHQRE
481 QRPVERKKPLIHYKEGMSPSEKPAWAKEIPHIVAVKSQGPALTASQSVHEQPTKGLSGFQE
541 ALNVTSHRVEMPRQNSDPTSENPLPTRIEKFDRSSWLRQEEIDIPPKVPPQRTTSISPALA
601 RKNSPGNGSALGPRLGSQPIRASNPDLRRTEPILESPLQRTSSGSSSSSTPSSQPSQG
661 GSQPGSQAGSSERTRVTRANSKSEGSPVLPHEPAVKPEESRDTIRPSRPAKYKKAIDEDL
721 TALAKELRELRIEETNRPMKKVTDYSSSEESSEEEEDGESETHDGTVAVSDIPRLI
781 PTGAPGSNEQYNVGMVGTGLETSHADSFSGSISREGTLMIRETSGEKRSGHSDSNGFA
841 GHINLPDLVQQSHSPACTPTEGLGRVSTHSQEMDSGTEYGMGSSTKASFTPFVDPRVYQT
901 SPTDEDEEDEESSAAALFTSELLRQEQQAKLNEARAKISVNVNNPNTNIRPHSDTPEIRKYKK
961 RFNSEILCAALWGVNLLVGTENGMLLDLRSQGQKVYNLINRRRFQQMDVILEGLNVLTIS
1021 GKKNKLRVYYLSQLRNRILHNDPEVEKQKGWITVGDLEGCIHYKVVKYERIKFLVIALKN
1081 AVEIYAWAPKPYHKFMAFKSFADLQHKLPLJLVDLTVEEQRLKVIFGSHTGFHVIDVDSGN
1141 SYDIYIPSHIQGNITPHAIIVILPKTDGEMMLVCYDEGVYVNTYGRITKDVLQWIGEMPT
1201 SVAYIHSNQIMGWGEKAIEIRSVEVGHLKFLCERNDKVFFASVRSGGS
1261 SQVFEMTLNRNSMMWZ

FIG.-32



1 MASDSPARSLDIISALRDPAGIFELVELVGNGTYGQVYKGRHVKTGQLAAIKVMDVTG
61 DEEEEIKQEINMLKKYSHHRNIAATYGAFAIKKKNNPPGMDDQLWLVMEMFCGAGSVTDLIKNT
121 KGNTLKEEWIAIYCIREILRGLSHLHQHKVIIHDIKGQNVLLTENAEVKLWDFGVSAQLDR
181 TVGRRNTFIGTPYUWMAPEVIACDENPDATYDFKSDLWSLIGITAIEMAEGAPPLCDMHPMR
241 ALFLIPRNPAPRLKSKWSKKFKQSFIESTCLVKNHSQRPATEQLMKHPFIRDQPNERQVR
301 QLKDHIDRTKPKRGEKDETEYEYSGSEEEEEENDSGEPSSILNLPGESTLRRDFLRLQLA
361 NKERSEALRRRQGLEQQQRNENEHKRQLLAERQKRRIEEQKEQRRRLEEQQRREKELRKQQE
421 REQRRHYEEQMRRREERRRAEHEQEYKRKQLEEQRQAERLQRQLKQERDYLVSLQHQQRQE
481 QRPVEKKPLYHYKEGMSPSEKPAWAKEVEERSRLNRQSSPAMPHKVANRISDPNLPPRSE
541 SFSISGVQPARTPPMRLRPVDPQIPIHLVAVKSQGPALTASQSVHEQPTKGILSGFQEA
601 SHRVEMPRQNSDPTSENPLPLPTRIEKFDRSSWLRLQEEIDIPPKVVPQRTTSISPALARKNSP
661 GNGSALGPRLGSQPIRASNPDLIRRTEPILSPLQRTSSGSSSSSTPSSQSSQGGSQPG
721 SQAGSSERTRVTRANSKSEGSVPVLPHEPAKVKPEESRDTITRPSRPAIDLTA
781 ETNRPMKKVTDYSSSEESSEEEEDGESETHDGTVAVSDIPRLIPTGAPGSNEQYNV
841 GMVGTGLETSHADSFSGSISREGTLMIRETSGEKKRSGHSDSNGFAGHINLPDLVQQSH
901 SPAGTPTEGLGRVSTHSQEMDSGTEYGMGSSTKASFTPFDPRVYQTSP
961 TDEDEEES
961 AAALFTSELLRQEQA
1021 VNLLVGTENGMLLDRSGQGKVVYNLINRRRFQQMDVLEGLNVLTISGKKNKLRVYYLSW
1081 LRNRILHNDPEVEKKQGWITVGDLLEGCIHYKVVKYERIKFLVIALKNAVEIYAWAPKPYH
1141 KFMAFKSFADLQHKPILLVDTVEEGQRLKVIFGSHTGFTVIDVDSGNSYDIYIPSHIQGN
1201 ITPHAIVILPKTDGMEMLVCYEDEGVYVNTYGRITKDVLQWGEMPTSVA
1261 GEKAIEIRSVEITGHLDGVFMKRAQRLLKFLCERN
1321 MNWZ

FIG.-33



1 MASDSPARSLDIEDLSDLRDPAGIFELVELVGNNGTYGQVYKGRHVKTGQLAAIKVMDVTG
61 DEEEEIKQEINMLKKYSHRNIAIATYYGAFIKKNPPGMDDQLWLWMEFCGAGSVTDLIKNT
121 KGNTLKEEWIAIYCIREILRGLSHLHQHKVIHRDIKGQNVLLENAAEVKLVDFFGVSQQLDR
181 TVGRRNTFIGTPYWMAPVIACDENPDATYDFEKSDLWLSIGITAJEMAEGAPPCLCDMHPMR
241 ALFLIPRNPNPAPRLKSKWSKFQSFIESCLVKNHSQRPAEQLMKHPFIRDQPNERQVR
301 QLKDHIDRTKKRGEIODETEYEYSGSEEEENDSGEPSSILNLPGESTLRRDFLRLQLA
361 NKERSEALRQQLEQQQRENEEHKRQLLAERQKRIEEQKERRQRRLEEQQRREKELRKQQE
421 REQRHYEEQMRREREERRRAEHEQEYIRRQLEEEQRQLEILQQQLHEQALLLEYKRKQL
481 EEQRQAERLQRQLKQERDYLVLSSLQHORQEQRPVKEKKPLYHYKEGMSPSEKPAWAKEIPH
541 VAVKSQGPALTASQSVHEQPTKGLSGFQEALNVTSHRVEMPRQNSDPTSENPPLEPTRIEK
601 FDRSSWLRQEEDIPPKVQPQRTTSISPALARKNSPGNGSALGPRLGSQPIRASNPDLRRTE
661 FILESPLQRTSSSSSSSTPSSQPSQGGSQPGSQAGSSERTVRANSKSEGSPVLPHE
721 PAKVKPEESRDITRPSRPADLTALAKELRELRIEETNRPMKKVTDYSSSEESSEEEE
781 EDGESETHDGTVAVDIPRPLIPTGAPGSNEQYNVGMVGHGLETSHADSFGSISREGTL
841 MIRETSGEKKRSGHSDSNGFAGHINLPDLVQQSHSPAGTPTEGLGRVSTHSQEMDSGTEY
901 GMGSSTKASFTPFDPRVYQTSPPTDEDEEDEESSAAALFTSELRLRQEQAQLNEARKISVV
961 NVNPTNIRPHSDTPEIRKYKKRFNSEIILCAALWGVNLJVGTTENGMLLDRSQGKVVNL
1021 NRRRFQMDVLEGNLVLTISGKKNKLRVYYLSWLNRNRLHNDPEVEKKQGWITVGDLEG
1081 CIHYKVKVVKYERIKFLVIALKNAVEIYAWAPKPYHKFMAFKSFAOLQHKPLLVDLTVEEGO
1141 RLKVIFGSHTGFHVIDVDSGNSYDIYIPSHIQGNITPHAIIVILPKTDGMEMLVCYEDEGV
1201 YVNTYGRITKDVLQWGEMPTSVAIHSNQIMGMGEKAIEIERSVETGHLDGVFMHKRAQR
1261 LKFLCERNDKVFASVRSGSSQVFFMTLNRNNSMMNWZ

FIG.-34



1 MASDSPARSLDIDLSALRDPAGIFELVELVNGNGTYGQVYKGRHVKTGQLAAIKVMDVTG
61 DEEEEIKQEINMLKKYSHHRNIAATYYGAFIKKNPPGMDDQLWLVMEFCGAGSVTDLIKNT
121 KGNTLKEEWIAYICREILRGLSHLHQHKVIRDIKGQNVLLTENAEVKLVDFGVSAQLDR
181 TVGRNNTFIGTPYWMAPEVIACDENPDATYDFKSDLWLSLIGITAIEEMAEGAPPLCDMHPMR
241 ALFLIPRNPNPAPRLKSKKWKSFFQSFIESCLVKNHSQRPAEQLMKHPFIRDQPNERQVR
301 QLKDHIDRTKKRGEKDETEYEYSGSEEEEEENDSGEPPSSILNLPGESTLRRDFLRLQLA
361 NKERSEALRRQQLEQQQRENEEHKRQLLAERQRKRIEEQKERRQRRLEEQQRREKEELRKQQE
421 REQRRHYEEQMRREEERRRAAEHEQEYKRKQLEEQRQAERLQRQLKQERDYLVSLQHQQRQE
481 QRPVEKKPLHYKEGMSPSEKPAWAKEIPHILVAVKSQGPALTASQSVHEQPTKGLSGFQE
541 ALNVTSHRVEMPRQNSDPTSENPPPLPTRIEKFDRSSWLQEEIDIPPKVVPQRTTSISPALA
601 RKNSPGNGSALGPRLGSQPIRASNPDLRRTEPILESPLQRTSSSSSSSTPSSQPSQSG
661 GSQPGSQAGSSERTVRTRANSKSEGSPVLPHEPAKVKPEESRDIITRPSRPAIDLTAIAKELR
721 ELRIETNRPMKKVTDYSSSSEEESESESSEEEEDGESETHDGTVAVSDIPLRPLIPTGAPGSN
781 EQYNVGMVGTHGLETSADSFSGSISREGTLMIRETSGEKKRSGHSDNSGFAGHINLPDL
841 VQQSHSPAGTPTEGLGRVSTHSQEMDSGTEYGMGSSTKASFTFPFDPRVYQTSPTDEEE
901 DEESSAAALFTSELLRQEAKLINEARKISVUVNNPTNIRPHSDTPEIRKYKRRFNSEILC
961 AALWGVNLLVGTENGMLLDRSGQGKVVYNLLINRRRFQQMDVLEGLNVLVTISGKKNNKLRV
1021 YYLSWLRNRILHNDPEVEKKQGWITVGDLEGCIHYKVKVYERIKFLVIALKNAVIEYAWA
1081 PKPYHKEMAFKSFADLQHKPLIQLVDTVEEGQRLKVIFGSHTGFFHVIDVDSGNSYDIYIPS
1141 HIQGNITPHAIIVILPKTDGMEMLVCYEDEGVYVNTYGRITKDVVLQWGEMPTSVAIHSN
1201 QIMGWGEKAIEIRSVEGTGHLDGVFMHKRAQRLKFLCERNDKVFASVRSGSSQVFEMTL
1261 NRNSMMNWZ

FIG.-35